CRF Errors Corrected by th STIC Systems Branch

CRF Processing Date: 6/17/2002

Serial Number: 10/010,667A

ENTERED

Edited by: _____

Verified by: _____ (STIC staff)

- [ ] Changed a file from non-ASCII to ASCII

- [ ] Changed the margins in cases where the sequence text was wrapped down to the next line.

- [ ] Edited a format error in the Current Application Data section, specifically: _____

- [ ] Edited the Current Application Data section with the actual current number. The number inputted by the applicant was [ ] the prior application data; or [ ] other _____.

- [ ] Added the mandatory heading and subheadings for "Current Application Data".

- [ ] Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

- [ ] Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

- [ ] Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

- [ ] Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____

- [ ] Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

- [ ] Inserted colons after headings/subheadings. Headings edited included: _____

- [ ] Deleted extra, invalid, headings used by an applicant, specifically: _____

- [✓] Deleted: [✓] non-ASCII "garbage" at the beginning/end of files; [ ] secretary initials/filename at end of file; [ ] page numbers throughout text; [ ] other invalid text, such as _____.

- [ ] Inserted mandatory headings, specifically: _____

- [ ] Corrected an obvious error in the response, specifically: _____

- [ ] Edited identifiers where upper case is used but lower case is required, or vice versa.

- [ ] Corrected an error in the Number of Sequences field, specifically: _____

- [ ] A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

- [ ] Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

- [ ] Other: _____
  _____
  _____
  _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

OIPE

RAW SEQUENCE LISTING   DATE: 06/17/2002
PATENT APPLICATION: US/10/010,667A   TIME: 19:48:45

Input Set : N:\jumbos\010667A.txt
Output Set: N:\CRF3\06172002\J010667A.raw

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 4 <110> APPLICANT: Afar, Daniel
 5         Hubert, Rene S.
 6         Leong, Kahan
 7         Raitano, Arthur B.
 8         Saffran, Douglas C.
 9         Mitchell, Steve Chappell
11 <120> TITLE OF INVENTION: NOVEL SERPENTINE TRANSMEMBRANE ANTIGENS
12         EXPRESSED IN HUMAN CANCERS AND USES THEREOF
15 <130> FILE REFERENCE: 511582001601
17 <140> CURRENT APPLICATION NUMBER: US 10/010,667A
18 <141> CURRENT FILING DATE: 2001-12-06
20 <150> PRIOR APPLICATION NUMBER: 09/323,873
21 <151> PRIOR FILING DATE: 1999-06-01
23 <150> PRIOR APPLICATION NUMBER: 60/087,520
24 <151> PRIOR FILING DATE: 1998-06-01
26 <150> PRIOR APPLICATION NUMBER: 60/091,183
27 <151> PRIOR FILING DATE: 1998-06-30
29 <160> NUMBER OF SEQ ID NOS: 32
31 <170> SOFTWARE: FastSEQ for Windows Version 4.0
33 <210> SEQ ID NO: 1
34 <211> LENGTH: 1195
35 <212> TYPE: DNA
36 <213> ORGANISM: Homo Sapiens
38 <400> SEQUENCE: 1
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40 aattaatgga aagcagaaaa gacatcacaa accaagaaga actttggaaa atgaagccta     120
41 ggagaaattt agaagaagac gattatttgc ataaggacac gggagagacc agcatgctaa     180
42 aaagacctgt gcttttgcat ttgcaccaaa cagcccatgc tgatgaattt gactgccctt     240
43 cagaacttca gcacacacag gaactctttc cacagtggca cttgccaatt aaaatagctg     300
44 ctattatagc atctctgact tttctttaca ctcttctgag ggaagtaatt caccctttag     360
45 caacttccca tcaacaatat ttttataaaa ttccaatcct ggtcatcaac aaagtcttgc     420
46 caatggtttc catcactctc ttggcattgg tttacctgcc aggtgtgata gcagcaattg     480
47 tccaacttca taatggaacc aagtataaga agtttccaca ttggttggat aagtggatgt     540
48 taacaagaaa gcagtttggg cttctcagtt tcttttttgc tgtactgcat gcaatttata     600
49 gtctgtctta cccaatgagg cgatcctaca gatacaagtt gctaaactgg gcatatcaac     660
50 aggtccaaca aaataaagaa gatgcctgga ttgagcatga tgtttggaga atggagattt     720
51 atgtgtctct gggaattgtg ggattggcaa tactggctct gttggctgtg acatctattc     780
52 catctgtgag tgactctttg acatggagag aatttcacta tattcagagc aagctaggaa     840
53 ttgtttccct tctactgggc acaatacacg cattgatttt tgcctggaat aagtggatag     900
54 atataaaaca atttgtatgg tatacacctc caacttttat gatagctgtt ttccttccaa     960
55 ttgttgtcct gatatttaaa agcatactat tcctgccatg cttgaggaag aagatactga    1020
56 agattagaca tggttgggaa gacgtcacca aaattaacaa aactgagata tgttcccagt    1080
57 tgtagaatta ctgtttacac acattttgt tcaatattga tatattttat caccaacatt    1140
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RAW SEQUENCE LISTING   DATE: 06/17/2002
PATENT APPLICATION: US/10/010,667A   TIME: 19:48:45

Input Set : N:\jumbos\010667A.txt
Output Set: N:\CRF3\06172002\J010667A.raw

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62 <212> TYPE: PRT
63 <213> ORGANISM: Homo sapiens
65 <400> SEQUENCE: 2
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67  1               5                  10                  15
68 Lys Pro Arg Arg Asn Leu Glu Glu Asp Asp Tyr Leu His Lys Asp Thr
69              20                  25                  30
70 Gly Glu Thr Ser Met Leu Lys Arg Pro Val Leu Leu His Leu His Gln
71          35                  40                  45
72 Thr Ala His Ala Asp Glu Phe Asp Cys Pro Ser Glu Leu Gln His Thr
73          50                  55                  60
74 Gln Glu Leu Phe Pro Gln Trp His Leu Pro Ile Lys Ile Ala Ala Ile
75 65                  70                  75                  80
76 Ile Ala Ser Leu Thr Phe Leu Tyr Thr Leu Leu Arg Glu Val Ile His
77              85                  90                  95
78 Pro Leu Ala Thr Ser His Gln Gln Tyr Phe Tyr Lys Ile Pro Ile Leu
79              100                 105                 110
80 Val Ile Asn Lys Val Leu Pro Met Val Ser Ile Thr Leu Leu Ala Leu
81          115                 120                 125
82 Val Tyr Leu Pro Gly Val Ile Ala Ala Ile Val Gln Leu His Asn Gly
83          130                 135                 140
84 Thr Lys Tyr Lys Lys Phe Pro His Trp Leu Asp Lys Trp Met Leu Thr
85 145                 150                 155                 160
86 Arg Lys Gln Phe Gly Leu Leu Ser Phe Phe Phe Ala Val Leu His Ala
87              165                 170                 175
88 Ile Tyr Ser Leu Ser Tyr Pro Met Arg Arg Ser Tyr Arg Tyr Lys Leu
89              180                 185                 190
90 Leu Asn Trp Ala Tyr Gln Gln Val Gln Gln Asn Lys Glu Asp Ala Trp
91          195                 200                 205
92 Ile Glu His Asp Val Trp Arg Met Glu Ile Tyr Val Ser Leu Gly Ile
93          210                 215                 220
94 Val Gly Leu Ala Ile Leu Ala Leu Leu Ala Val Thr Ser Ile Pro Ser
95 225                 230                 235                 240
96 Val Ser Asp Ser Leu Thr Trp Arg Glu Phe His Tyr Ile Gln Ser Lys
97              245                 250                 255
98 Leu Gly Ile Val Ser Leu Leu Leu Gly Thr Ile His Ala Leu Ile Phe
99              260                 265                 270
100 Ala Trp Asn Lys Trp Ile Asp Ile Lys Gln Phe Val Trp Tyr Thr Pro
101          275                 280                 285
102 Pro Thr Phe Met Ile Ala Val Phe Leu Pro Ile Val Val Leu Ile Phe
103          290                 295                 300
104 Lys Ser Ile Leu Phe Leu Pro Cys Leu Arg Lys Lys Ile Leu Lys Ile
105 305                 310                 315                 320
106 Arg His Gly Trp Glu Asp Val Thr Lys Ile Asn Lys Thr Glu Ile Cys
107              325                 330                 335
108 Ser Gln Leu
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113 <212> TYPE: DNA
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116 <400> SEQUENCE: 3
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121 <211> LENGTH: 24
122 <212> TYPE: DNA
123 <213> ORGANISM: Artificial Sequence
125 <220> FEATURE:
126 <223> OTHER INFORMATION: Primer
128 <400> SEQUENCE: 4
129 actttgttga tgaccaggat tgga                                           24
131 <210> SEQ ID NO: 5
132 <211> LENGTH: 24
133 <212> TYPE: DNA
134 <213> ORGANISM: Artificial Sequence
136 <220> FEATURE:
137 <223> OTHER INFORMATION: Primer
139 <400> SEQUENCE: 5
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143 <211> LENGTH: 3627
144 <212> TYPE: DNA
145 <213> ORGANISM: Homo sapiens
147 <400> SEQUENCE: 6
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149 gtgggtggct gaagccatac tattttatag aattaatgga aagcagaaaa gacatcacaa    120
150 accaagaaga actttggaaa atgaagccta ggagaaattt agaagaagac gattatttgc    180
151 ataaggacac gggagagacc agcatgctaa aaagacctgt gctttttgcat ttgcaccaaa   240
152 cagcccatgc tgatgaattt gactgccctt cagaacttca gcacacacag gaactctttc    300
153 cacagtggca cttgccaatt aaaatagctg ctattatagc atctctgact tttctttaca    360
154 ctcttctgag ggaagtaatt caccccttag caacttccca tcaacaatat ttttataaaa    420
155 ttccaatcct ggtcatcaac aaagtcttgc caatggtttc catcactctc ttggcattgg    480
156 tttacctgcc aggtgtgata gcagcaattg tccaacttca taatggaacc aagtataaga    540
157 agtttccaca ttggttggat aagtggatgt taacaagaaa gcagtttggg cttctcagtt    600
158 tcttttttgc tgtactgcat gcaatttata gtctgtctta cccaatgagg cgatcctaca    660
159 gatacaagtt gctaaactgg gcatatcaac aggtccaaca aaataaagaa gatgcctgga    720
160 ttgagcatga tgtttggaga atggagattt atgtgtctct gggaattgtg ggattggcaa    780
161 tactggctct gttggctgtg acatctattc catctgtgag tgactctttg acatggagag    840
162 aatttcacta tattcaggta aataatatat aaaataaccc taagaggtaa atcttctttt    900
163 tgtgtttatg atatagaata tgttgacttt accccataaa aaataacaaa tgttttttcaa   960
164 cagcaaagat cttatacttg ttccaattaa taatgtgctc tcctgttgtt ttccctattg   1020
165 cttctaatta ggacaagtgt ttcctagaca taaataaaag gcattaaaat attctttgtt   1080
166 tttttttttt tgtttgtttg tttttttgttt gtttgtttgt tttttttgaga tgaagtctcg   1140
167 ctctgttgcc catgctggag tacagtggca cgatctcggc tcactgcaac ctgcgcctcc   1200
168 tgggttcagg cgattctctt gcctcagcct cctgagtagc tgggattaca ggcacccatc   1260
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169 accatgtcca gctaattttt gtatttttag tagagacagg gttttcccat gttggccagg    1320
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171 tgacagttgt gagccaccac actcagcctg ctctttctaa tatttgaaac ttgttagaca    1440
172 atttgctacc catctaatgt gatattttag gaatccaata tgcatggttt attatttctt    1500
173 aaaaaaaata ttcttttacc tgtcacctga atttagtaat gccttttatg ttacacaact    1560
174 tagcactttc cagaaacaaa aactctctcc ttgaaataat agagttttta tctaccaaag    1620
175 atatgctagt gtctcatttc aaaggctgct ttttccagct tacatttttat atacttactc    1680
176 acttgaagtt tctaaatatt cttgtaattt taaaactatc tcagatttac tgaggtttat    1740
177 cttctggtgg tagattatcc ataagaagag tgatgtgcca gaatcactct gggatccttg    1800
178 tctgacaaga ttcaaaggac taaatttaat tcagtcatga acactgccaa ttaccgttta    1860
179 tgggtagaca tctttggaaa tttccacaag gtcagacatt cgcaactatc ccttctacat    1920
180 gtccacacgt atactccaac actttattag gcatctgatt agtttggaaa gtatgcctcc    1980
181 atctgaatta gtccagtgtg gcttagagtt ggtacaacat tctcacagaa tttcctaatt    2040
182 ttgtaggttc agcctgataa ccactggagt tctttggtcc tcattaaata gctttcttca    2100
183 cacattgctc tgcctgttac acatatgatg aacactgctt tttagacttc attaggaatt    2160
184 taggactgca tcttgacaac tgagcctatt ctactatatg tacaatacct agcccataat    2220
185 aggtatacaa tacacatttg gtaaaactaa ttttcaacca atgacatgta tttttcaact    2280
186 agtaacctag aaatgtttca cttaaaatct gagaactggt tacactacaa gttaccttgg    2340
187 agattcatat atgaaaacgc aaacttagct atttgattgt attcactggg acttaagaat    2400
188 gcgcctgaat aattgtgagt tcgatttgtt ctggcaggct aatgaccatt tccagtaaag    2460
189 tgaatagagg tcagaagtcg tataaaagag gtgttgtcag aacaccgttg agattacata    2520
190 ggtgaacaac tattttttaag caactttatt tgtgtagtga caaagcatcc caatgcaggc    2580
191 tgaaatgttt catcacatct ctggatctct ctattttgtg cagacattga aaaaattgtt    2640
192 catattattt ccatgttatc agaatatttg attttttaaa aacataggcc aagttcattc    2700
193 acttcattat tcatttatca aaatcagagt gaatcacatt agtcgccttc acaactgata    2760
194 aagatcactg aagtcaaatt gattttttgct ataatcttca atctacctat atttaattga    2820
195 gaatctaaaa tgtacaaatc attgtgttga ttctgcagtg atcctgctat aagtaagact    2880
196 cagtccctga tttttaggtat cctgtgaaaa gcagaattaa gacaaataca .caagagacaa    2940
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198 tttttgatca cctgccttca aagaaaggct gtgaattttg ttcacttaga cagcttggag    3060
199 acaagaaatt acccaaaagt aaggtgaggg ggataggcaa aaagagcaga aagatgtgaa    3120
200 tggacattgt tgagaaatgt gataggaaaa caatcataga taaaggattt ccaagcaaca    3180
201 gagcatatcc agatgaggta ggatgggata aactcttatt gaaccaatct tcaccaattt    3240
202 tgttttttctt ttgcagagca agctaggaat tgtttccctt ctactgggca caatacacgc    3300
203 attgattttt gcctggaata agtggataga tataaaacaa tttgtatggt atacacctcc    3360
204 aacttttata atagctgttt tccttccaat tgttgtcctg atatttaaaa gcatactatt    3420
205 cctgccatgc ttgaggaaga agatactgaa gattagacat ggttgggaag acgtcaccaa    3480
206 aattaacaaa actgagatat gttcccagtt gtagaattac tgtttacaca cattttttgtt    3540
207 caatattgat atattttatc accaacattt caagtttgta tttgttaata aaatgattat    3600
208 tcaaggaaaa aaaaaaaaaa aaaaaaa    3627
210 <210> SEQ ID NO: 7
211 <211> LENGTH: 519
212 <212> TYPE: DNA
213 <213> ORGANISM: Homo sapiens
215 <400> SEQUENCE: 7
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217 ttgctctccc tagtatacct cgcaggtctt ctggcagctg cttatcaact ttattacggc    120
218 accaagtata ggagatttcc accttggttg gaaacctggt tacagtgtag aaaacagctt    180
219 ggattactaa gttttttctt cgctatggtc catgttgcct acagcctctg cttaccgatg    240
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                    RAW SEQUENCE LISTING                  DATE: 06/17/2002
                    PATENT APPLICATION:  US/10/010,667A    TIME: 19:48:45

                    Input Set : N:\jumbos\010667A.txt
                    Output Set: N:\CRF3\06172002\J010667A.raw

220 agaaggtcag agagatattt gtttctcaac atggcttatc agcaggttca tgcaaatatt    300
221 gaaaactctt ggaatgagga agaagtttgg agaattgaaa tgtatatctc ctttggcata    360
222 atgagccttg gcttactttc cctcctggca gtcacttcta tcccttcagt gagcaatgct    420
223 ttaaactgga gagaattcag ttttattcag tctacacttg gatatgtcgc tctgctcata    480
224 agtactttcc atgttttaat ttatggatgg aaacgagct                           519
226 <210> SEQ ID NO: 8
227 <211> LENGTH: 173
228 <212> TYPE: PRT
229 <213> ORGANISM: Homo sapiens
231 <400> SEQUENCE: 8
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233 1               5                   10                  15
234 Val Ala Ile Thr Leu Leu Ser Leu Val Tyr Leu Ala Gly Leu Leu Ala
235     20                  25                  30
236 Ala Ala Tyr Gln Leu Tyr Tyr Gly Thr Lys Tyr Arg Arg Phe Pro Pro
237     35                  40                  45
238 Trp Leu Glu Thr Trp Leu Gln Cys Arg Lys Gln Leu Gly Leu Leu Ser
239 50                  55                  60
240 Phe Phe Phe Ala Met Val His Val Ala Tyr Ser Leu Cys Leu Pro Met
241 65                  70                  75                  80
242 Arg Arg Ser Glu Arg Tyr Leu Phe Leu Asn Met Ala Tyr Gln Gln Val
243             85                  90                  95
244 His Ala Asn Ile Glu Asn Ser Trp Asn Glu Glu Glu Val Trp Arg Ile
245         100                 105                 110
246 Glu Met Tyr Ile Ser Phe Gly Ile Met Ser Leu Gly Leu Leu Ser Leu
247         115                 120                 125
248 Leu Ala Val Thr Ser Ile Pro Ser Val Ser Asn Ala Leu Asn Trp Arg
249     130                 135                 140
250 Glu Phe Ser Phe Ile Gln Ser Thr Leu Gly Tyr Val Ala Leu Leu Ile
251 145                 150                 155                 160
252 Ser Thr Phe His Val Leu Ile Tyr Gly Trp Lys Arg Ala
253             165                 170
255 <210> SEQ ID NO: 9
256 <211> LENGTH: 322
257 <212> TYPE: DNA
258 <213> ORGANISM: Homo sapiens
260 <400> SEQUENCE: 9
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262 cagagtgact tttacaaaat tcctatagag attgtgaata aaacttacc tatagttgcc     120
263 attactttgc tctccctagt ataccttgca ggtcttctgg cagctgctta tcaactttat    180
264 tacggcacca agtataggag atttccacct tggttggaaa cctggttaca gtgtagaaaa    240
265 cagcttggat tactaagttg tttcttcgct atggtccatg ttgcctacag cctctgctta    300
266 ccgatgagaa ggtcagagag at                                             322
268 <210> SEQ ID NO: 10
269 <211> LENGTH: 183
270 <212> TYPE: DNA
271 <213> ORGANISM: Homo sapiens
273 <400> SEQUENCE: 10
274 tttgcagctt tgcagatacc cagactgagc tggaactgga atttgtcttc ctattgactc    60
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 06/17/2002
PATENT APPLICATION: US/10/010,667A   TIME: 19:48:46

Input Set : N:\jumbos\010667A.txt
Output Set: N:\CRF3\06172002\J010667A.raw

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:12; N Pos. 11,56,233,250,310,326,377,398